



## SECOND SUBSTITUTE SEQUENCE LISTING

<110> Balint, Robert F.  
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KaloBios, Inc.

<120> Interaction-Activated Proteins

<130> 021167-000700US

<140> US 09/526,106

<141> 2000-03-15

<150> US 60/124,339

<151> 1999-03-15

<150> US 60/135,926

<151> 1999-05-25

<150> US 60/175,968

<151> 2000-01-13

<160> 27

<170> PatentIn Ver. 2.1

<210> 1

<211> 789

<212> DNA

<213> Escherichia coli

<220>

<221> CDS

<222> (1)..(789)

<223> TEM-1 beta-lactamase

<400> 1

cac cca gaa acg ctg gtg aaa gta aaa gat gct gaa gat cag ttg ggt 48  
His Pro Glu Thr Leu Val Lys Val Lys Asp Ala Glu Asp Gln Leu Gly  
1 5 10 15

gca cga gtg ggt tac atc gaa ctg gat ctc aac agc ggt aag atc ctt 96  
Ala Arg Val Gly Tyr Ile Glu Leu Asp Leu Asn Ser Gly Lys Ile Leu  
20 25 30

gag agt ttt cgc ccc gaa gaa cgt ttt cca atg atg agc act ttt aaa 144

Glu Ser Phe Arg Pro Glu Glu Arg Phe Pro Met Met Ser Thr Phe Lys  
35 40 45

gtt ctg cta tgt ggc gcg gta tta tcc cgt att gac gcc ggg caa gag 192  
Val Leu Leu Cys Gly Ala Val Leu Ser Arg Ile Asp Ala Gly Gln Glu  
50 55 60

caa ctc ggt cgc cgc ata cac tat tct cag aat gac ttg gtt gag tac 240  
Gln Leu Gly Arg Arg Ile His Tyr Ser Gln Asn Asp Leu Val Glu Tyr  
65 70 75 80

tca cca gtc aca gaa aag cat ctt acg gat ggc atg aca gta aga gaa 288  
Ser Pro Val Thr Glu Lys His Leu Thr Asp Gly Met Thr Val Arg Glu  
85 90 95

tta tgc agt get gcc ata acc atg agt gat aac act gcg gcc aac tta 336  
Leu Cys Ser Ala Ala Ile Thr Met Ser Asp Asn Thr Ala Ala Asn Leu  
100 105 110

ctt ctg aca acg atc gga gga ccg aag gag cta acc gct ttt ttg cac 384  
Leu Leu Thr Thr Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe Leu His  
115 120 125

aac atg ggg gat cat gta act cgc ctt gat cgt tgg gaa ccg gag ctg 432  
Asn Met Gly Asp His Val Thr Arg Leu Asp Arg Trp Glu Pro Glu Leu  
130 135 140

aat gaa gcc ata cca aac gac gag cgt gac acc acg atg cct gta gca 480  
Asn Glu Ala Ile Pro Asn Asp Glu Arg Asp Thr Thr Met Pro Val Ala  
145 150 155 160

atg gca aca acg ttg cgc aaa cta tta act ggc gaa cta ctt act cta 528  
Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu Thr Leu  
165 170 175

gct tcc cgg caa caa tta ata gac tgg atg gag gcg gat aaa gtt gca 576  
Ala Ser Arg Gln Gln Leu Ile Asp Trp Met Glu Ala Asp Lys Val Ala  
180 185 190

gga cca ctt ctg cgc tcg gcc ctt ccg gct ggc tgg ttt att gct gat 624  
Gly Pro Leu Leu Arg Ser Ala Leu Pro Ala Gly Trp Phe Ile Ala Asp  
195 200 205

aaa tct gga gcc ggt gag cgt ggg tct cgc ggt atc att gca gca ctg 672  
Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala Ala Leu  
210 215 220

ggg cca gat ggt aag ccc tcc cgt atc gta gtt atc tac acg acg ggg 720  
 Gly Pro Asp Gly Lys Pro Ser Arg Ile Val Val Ile Tyr Thr Thr Gly  
 225 230 235 240

agt cag gca act atg gat gaa cga aat aga cag atc gct gag ata ggt 768  
 Ser Gln Ala Thr Met Asp Glu Arg Asn Arg Gln Ile Ala Glu Ile Gly  
 245 250 255

gcc tca ctg att aag cat tgg 789  
 Ala Ser Leu Ile Lys His Trp  
 260

<210> 2

<211> 263

<212> PRT

<213> Escherichia coli

<220>

<223> TEM-1 beta-lactamase

<400> 2

His Pro Glu Thr Leu Val Lys Val Lys Asp Ala Glu Asp Gln Leu Gly  
 1 5 10 15

Ala Arg Val Gly Tyr Ile Glu Leu Asp Leu Asn Ser Gly Lys Ile Leu  
 20 25 30

Glu Ser Phe Arg Pro Glu Glu Arg Phe Pro Met Met Ser Thr Phe Lys  
 35 40 45

Val Leu Leu Cys Gly Ala Val Leu Ser Arg Ile Asp Ala Gly Gln Glu  
 50 55 60

Gln Leu Gly Arg Arg Ile His Tyr Ser Gln Asn Asp Leu Val Glu Tyr  
 65 70 75 80

Ser Pro Val Thr Glu Lys His Leu Thr Asp Gly Met Thr Val Arg Glu  
 85 90 95

Leu Cys Ser Ala Ala Ile Thr Met Ser Asp Asn Thr Ala Ala Asn Leu  
 100 105 110

Leu Leu Thr Thr Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe Leu His  
 115 120 125

Asn Met Gly Asp His Val Thr Arg Leu Asp Arg Trp Glu Pro Glu Leu  
130 135 140

Asn Glu Ala Ile Pro Asn Asp Glu Arg Asp Thr Thr Met Pro Val Ala  
145 150 155 160

Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu Thr Leu  
165 170 175

Ala Ser Arg Gln Gln Leu Ile Asp Trp Met Glu Ala Asp Lys Val Ala  
180 185 190

Gly Pro Leu Leu Arg Ser Ala Leu Pro Ala Gly Trp Phe Ile Ala Asp  
195 200 205

Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala Ala Leu  
210 215 220

Gly Pro Asp Gly Lys Pro Ser Arg Ile Val Val Ile Tyr Thr Thr Gly  
225 230 235 240

Ser Gln Ala Thr Met Asp Glu Arg Asn Arg Gln Ile Ala Glu Ile Gly  
245 250 255

Ala Ser Leu Ile Lys His Trp  
260

<210> 3

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:linker

<400> 3

Gly Gly Gly Gly Ser

1

5

<210> 4

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:flexible linker

<400> 4

Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser  
1 5 10 15

<210> 5

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:hexa-histidine  
tag

<400> 5

His His His His His His  
1 5

<210> 6

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:flexible linker  
of variable length

<220>

<221> REPEAT

<222> (1)..(5)

<223> (G-4S)-x, amino acids 1-5 may be repeated an  
undefined number of times

<400> 6

Gly Gly Gly Gly Ser  
1 5

<210> 7

<211> 267

<212> PRT

<213> Escherichia coli

<220>

<223> Neomycin phosphotransferase II (NPTII)

<400> 7

Met Gly Ser Ala Ile Glu Gln Asp Gly Leu His Ala Gly Ser Pro Ala  
1 5 10 15

Ala Trp Val Glu Arg Leu Phe Gly Tyr Asp Trp Ala Gln Gln Thr Ile  
20 25 30

Gly Cys Ser Asp Ala Ala Val Phe Arg Leu Ser Ala Gln Gly Arg Pro  
35 40 45

Val Leu Phe Val Lys Thr Asp Leu Ser Gly Ala Leu Asn Glu Leu Gln  
50 55 60

Asp Glu Ala Ala Arg Leu Ser Trp Leu Ala Thr Thr Gly Val Pro Cys  
65 70 75 80

Ala Ala Val Leu Asp Val Val Thr Glu Ala Gly Arg Asp Trp Leu Leu  
85 90 95

Leu Gly Glu Val Pro Gly Gln Asp Leu Leu Ser Ser His Leu Ala Pro  
100 105 110

Ala Glu Lys Val Ser Ile Met Ala Asp Ala Met Arg Arg Leu His Thr  
115 120 125

Leu Asp Pro Ala Thr Cys Pro Phe Asp His Gln Ala Lys His Arg Ile  
130 135 140

Glu Arg Ala Arg Thr Arg Met Glu Ala Gly Leu Val Asp Gln Asp Asp  
145 150 155

Leu Asp Glu Glu His Gln Gly Leu Ala Pro Ala Glu Leu Phe Ala Arg  
160 165 170 175

Leu Lys Ala Arg Met Pro Asp Gly Glu Asp Leu Val Val Thr His Gly  
180 185 190

Asp Ala Cys Leu Pro Asn Ile Met Val Glu Asn Gly Arg Phe Ser Gly  
195 200 205

Phe Ile Asp Cys Gly Arg Leu Gly Val Ala Asp Arg Tyr Gln Asp Ile

210	215	220
Ala Leu Ala Thr Arg Asp Ile Ala Glu Glu Leu Gly Gly Glu Trp Ala		
225	230	235
Asp Arg Phe Leu Val Leu Tyr Gly Ile Ala Ala Pro Asp Ser Gln Arg		
240	245	250 255
Ile Ala Phe Tyr Arg Leu Leu Asp Glu Phe Phe		
260	265	

<210> 8  
 <211> 18  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence:CD40-binding  
 Trxpep

<400> 8  
 Cys Gly Pro Lys Glu Leu Arg Ile Gly Gly Arg Pro Arg Arg Pro Gly  
 1 5 10 15

Pro Cys

<210> 9  
 <211> 18  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence:CD40-binding  
 Trxpep

<400> 9  
 Cys Gly Pro Glu Gly Gln Gly Gly Val Ala Val Gly Gly Val Gly Gly  
 1 5 10 15

Pro Cys

<210> 10  
 <211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 10

Cys Gly Pro Ala Lys Arg Ala Asp Val Glu Phe Ser Leu Glu Pro Gly  
1 5 10 15

<210> 11

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 11

Ala Lys Pro Cys Gly Gln Gln Ser Ile His Leu Gly Gly Val Phe Glu  
1 5 10 15

Leu Gln Pro Gly Ala  
20

<210> 12

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 12

Cys Gly Pro Lys Ser Ala Gly Lys Gly Arg Lys Asp Arg Arg Lys Gly  
1 5 10 15

Pro Cys

<210> 13



<211> 19  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 13  
Cys Gly Pro Pro Arg Thr Arg Val Asn His Gln Gly Gln Lys Thr Arg  
1 5 10 15

Gly Pro Cys

<210> 14  
<211> 18  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 14  
Cys Gly Pro Ala Gly Ala Ile Arg His Glu His Arg Gln Gly Leu Gly  
1 5 10 15

Pro Cys

<210> 15  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 15  
Leu Val Thr Leu Glu Asn Gly Lys Gln Leu Thr Val Lys Arg Gln Gly  
1 5 10 15

Leu Tyr Tyr Ile Tyr Ala Gln  
20

<210> 16  
<211> 18  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 16  
Cys Gly Pro Asp Thr Gly Leu Glu Thr Asp Ala Ala Asp Ala Ser Gly  
1 5 10 15

Pro Cys

<210> 17  
<211> 18  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 17  
Cys Gly Pro Arg Arg Val Arg Glu Thr Val Ala Val Glu Ser Ser Gly  
1 5 10 15

Pro Cys

<210> 18  
<211> 18  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 18  
Cys Gly Pro Pro Cys Ala Thr Phe Glu Glu Ala Lys Ser Asn Gln Gly  
1 5 10 15

Pro Cys

<210> 19

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 19

Glu Thr Lys Lys Glu Asn Ser Phe Glu Met Gln Lys Gly Asp Gln Asn  
1 5 10 15

Pro Gln

<210> 20

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 20

Cys Gly Pro Gly Arg Glu Ser Arg Gly Arg Cys Tyr Thr Pro Ser Gly  
1 5 10 15

Pro Cys

<210> 21

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 21

Thr Asp Pro Ser Gln Val Ser His Gly Thr Gly Phe Thr Ser Phe Gly  
1 5 10 15

Leu Leu

<210> 22

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 22

Cys Gly Pro Asn Thr Pro Asp Glu Glu Met Ala Pro Gln Ala Pro Gly  
1 5 10 15

Pro Cys

<210> 23

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding  
Trxpep

<400> 23

Cys Gly Pro Val Val His Ile Lys Thr Asn Glu Gln Ala Ala Pro Gly  
1 5 10 15

Pro Cys

<210> 24

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:CD40-binding

Trxpep

<400> 24

Cys Gly Pro Val Ala Glu Glu Pro Ala Gly Gly Ala Gly Arg Pro Gly  
1                      5                      10                      15

Pro Cys

<210> 25

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Her-2/neu  
Tyr1068 phosphorylation substrate peptide

<400> 25

Pro Val Pro Glu Tyr Ile Asn Gln Ser  
1                      5

<210> 26

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:short flexible  
linker

<400> 26

Pro Gly Ser Gly Gly  
1                      5

<210> 27

<211> 263

<212> PRT

<213> Escherichia coli

<220>

<223> N-terminal beta-lactamase fragment

<400> 2

His Pro Glu Thr Leu Val Lys Val Lys Asp Ala Glu Asp Gln Leu Gly  
1 5 10 15

Ala Arg Val Gly Tyr Ile Glu Leu Asp Leu Asn Ser Gly Lys Ile Leu  
20 25 30

Glu Ser Phe Arg Pro Glu Glu Arg Phe Pro Met Met Ser Thr Phe Lys  
35 40 45

Val Leu Leu Cys Gly Ala Val Leu Ser Arg Ile Asp Ala Gly Gln Glu  
50 55 60

Gln Leu Gly Arg Arg Ile His Tyr Ser Gln Asn Asp Leu Val Glu Tyr  
65 70 75 80

Ser Pro Val Thr Glu Lys His Leu Thr Asp Gly Met Thr Val Arg Glu  
85 90 95

Leu Cys Ser Ala Ala Ile Thr Met Ser Asp Asn Thr Ala Ala Asn Leu  
100 105 110

Leu Leu Thr Thr Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe Leu His  
115 120 125

Asn Met Gly Asp His Val Thr Arg Leu Asp Arg Trp Glu Pro Glu Leu  
130 135 140

Asn Glu Ala Ile Pro Asn Asp Glu Arg Asp Thr Thr Thr Pro Val Ala  
145 150 155 160

Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu Thr Leu  
165 170 175

Ala Ser Arg Gln Gln Leu Ile Asp Trp Met Glu Ala Asp Lys Val Ala  
180 185 190

Gly Pro Leu Leu Arg Ser Ala Leu Pro Ala Gly Trp Phe Ile Ala Asp  
195 200 205

Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala Ala Leu  
210 215 220

Gly Pro Asp Gly Lys Pro Ser Arg Ile Val Val Ile Tyr Thr Thr Gly  
225 230 235 240

Ser Gln Ala Thr Met Asp Glu Arg Asn Arg Gln Ile Ala Glu Ile Gly

245

250

255

Ala Ser Leu Ile Lys His Trp  
260

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